JA PCT/09

CRF Errors Corrected by the STIC System Branch

Serial Number: 09/868,338

CRF Processing Date: 2/26/2002

Edited by:

Verified by: (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically:

  _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:

  _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

  _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

  _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included:

  _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically:

  _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____.
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically:

  _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically:

  _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☑ Other: Seq 7 - inserted hard return

*Examiner: The above corrections must be communicated to th applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

PCT09

RAW SEQUENCE LISTING DATE: 02/26/2002
PATENT APPLICATION: US/09/868,338 TIME: 12:46:49

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Output Set: N:\CRF3\02262002\I868338.raw

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  2       MATSUI, KAZUHIKO
  3       NAKAMATSU, TSUYOSHI
  4       KIMURA, EIICHIRO
  5 <120> TITLE OF INVENTION: ABC TRANSPORTER AND GENE CODING FOR THE SAME
  6 <130> FILE REFERENCE: 209861US-8222-10-0-PCT
  7 <140> CURRENT APPLICATION NUMBER: US/09/868,338
  8 <141> CURRENT FILING DATE: 2001-06-18
  9 <150> PRIOR APPLICATION NUMBER: JP 10/360621
 10 <151> PRIOR FILING DATE: 1998-12-18
 11 <150> PRIOR APPLICATION NUMBER: PCT/JP 99/07079
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 44 <222> LOCATION: (4)..(4)
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RAW SEQUENCE LISTING DATE: 02/26/2002
PATENT APPLICATION: US/09/868,338 TIME: 12:46:49

Input Set : N:\Crf3\02132002\I868338.raw
Output Set: N:\CRF3\02262002\I868338.raw

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RAW SEQUENCE LISTING DATE: 02/26/2002
PATENT APPLICATION: US/09/868,338 TIME: 12:46:49

Input Set : N:\Crf3\02132002\I868338.raw
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105 1 5 10 15
106 gga tcg cta acc ggt aac ctc agt gaa cta cgt gca caa act act ttt 96
107 Gly Ser Leu Thr Gly Asn Leu Ser Glu Leu Arg Ala Gln Thr Thr Phe
108 20 25 30
109 agt aca tta tgg gat acc cat aaa gaa acc tat aga gtc tcc ata gct 144
110 Ser Thr Leu Trp Asp Thr His Lys Glu Thr Tyr Arg Val Ser Ile Ala
111 35 40 45
112 tcc gca gca gga caa gac ttc tac ggg ctt gct gag act cta cgc act 192
113 Ser Ala Ala Gly Gln Asp Phe Tyr Gly Leu Ala Glu Thr Leu Arg Thr
114 50 55 60
115 atg gat agg cat ggg gaa att att ttg gca gat cgt caa tgg tta aca 240
116 Met Asp Arg His Gly Glu Ile Ile Leu Ala Asp Arg Gln Trp Leu Thr
117 65 70 75 80
118 gct ccc ctt gat atc ggt gca cca gtc gta tta tca aac aca act ttt 288
119 Ala Pro Leu Asp Ile Gly Ala Pro Val Val Leu Ser Asn Thr Thr Phe
120 85 90 95
121 gcc gtt gat gaa gga cta ctt gcg cca aaa gat cta ccg caa agt gac 336
122 Ala Val Asp Glu Gly Leu Leu Ala Pro Lys Asp Leu Pro Gln Ser Asp
123 100 105 110
124 gag atc aca ata ttg cat cct cag ttt ctg gat tcg gcc aaa gag cca 384
125 Glu Ile Thr Ile Leu His Pro Gln Phe Leu Asp Ser Ala Lys Glu Pro
126 115 120 125
127 gaa tta ctt ggt ttg ctg gag ttc gaa gca tcc aac tca caa gtg cca 432
128 Glu Leu Leu Gly Leu Leu Glu Phe Glu Ala Ser Asn Ser Gln Val Pro
129 130 135 140
130 atg cca aag atc caa agc att cca tat gat agc gaa gac tca acc aac 480
131 Met Pro Lys Ile Gln Ser Ile Pro Tyr Asp Ser Glu Asp Ser Thr Asn
132 145 150 155 160
133 ccc atg tct gaa gtt ttt acc tac aac att aac ctg gat agt gca gta 528
134 Pro Met Ser Glu Val Phe Thr Tyr Asn Ile Asn Leu Asp Ser Ala Val
135 165 170 175
136 aga aac cca atc gta gtt atc ctt ccc gca ggc tta gag ctt tta agt 576
137 Arg Asn Pro Ile Val Val Ile Leu Pro Ala Gly Leu Glu Leu Leu Ser
138 180 185 190
139 gat caa aat ttg tcg gct cga ctc aca cag aat agt ctg ctg ata aaa 624
140 Asp Gln Asn Leu Ser Ala Arg Leu Thr Gln Asn Ser Leu Leu Ile Lys
141 195 200 205
142 gac cag act ggt gtg aac gct ctt cta tcc tca gag gat tca cgc aat 672
143 Asp Gln Thr Gly Val Asn Ala Leu Leu Ser Ser Glu Asp Ser Arg Asn
144 210 215 220
145 tat gtg gga gct gca tcc ccg atg att gac acg tgg gaa gaa agc gtt 720
146 Tyr Val Gly Ala Ala Ser Pro Met Ile Asp Thr Trp Glu Glu Ser Val
147 225 230 235 240
148 gtt cgg ttg aag gaa gcg aac caa ata atc gcc ttc aac gct ttc att 768

RAW SEQUENCE LISTING DATE: 02/26/2002
PATENT APPLICATION: US/09/868,338 TIME: 12:46:49

Input Set : N:\Crf3\02132002\I868338.raw
Output Set: N:\CRF3\02262002\I868338.raw

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150                       245                 250                 255
151       gca ttg ttc ctc acg acg act ctt gtt cta gca tac tgc act ggt att      816
152       Ala Leu Phe Leu Thr Thr Thr Leu Val Leu Ala Tyr Cys Thr Gly Ile
153                   260                 265                 270
154       tca ttt aag aaa tca aag aag act atg ggt agc gca tct act agg aaa      864
155       Ser Phe Lys Lys Ser Lys Lys Thr Met Gly Ser Ala Ser Thr Arg Lys
156               275                 280                 285
157       tca tcc att aag agc tcg att aca gct gct aat tgt aga agt aat ttt      912
158       Ser Ser Ile Lys Ser Ser Ile Thr Ala Ala Asn Cys Arg Ser Asn Phe
159           290                 295                 300
160       cgc ttc aat tcc gtg cgt ctg gct cgc gaa ccg cta ttt cga gcg atc      960
161       Arg Phe Asn Ser Val Arg Leu Ala Arg Glu Pro Leu Phe Arg Ala Ile
162       305                 310                 315                 320
163       tgc agc aat agc ttc aga tgc tcc ctc agc cag ata ctt aga aca tct     1008
164       Cys Ser Asn Ser Phe Arg Cys Ser Leu Ser Gln Ile Leu Arg Thr Ser
165                       325                 330                 335
166       caa ttc tat acc tcc atc act gcc gtt ggt ttt agg aat ctt aat aat     1056
167       Gln Phe Tyr Thr Ser Ile Thr Ala Val Gly Phe Arg Asn Leu Asn Asn
168                   340                 345                 350
169       cgg ttg gac ttc act ttc att ttt cag ttc gat gaa gct tcc ttt         1101
170       Arg Leu Asp Phe Thr Phe Ile Phe Gln Phe Asp Glu Ala Ser Phe
171               355                 360                 365
172       tgaaaagagc acaca atg ata gaa atc aat gac ctc aag aaa tct ttt ggc     1152
173                        Met Ile Glu Ile Asn Asp Leu Lys Lys Ser Phe Gly
174                             370                 375
175       gtt cgg atc tta tgg caa ggt ctc agt cat aag ttt tta cca gga aca      1200
176       Val Arg Ile Leu Trp Gln Gly Leu Ser His Lys Phe Leu Pro Gly Thr
177       380                 385                 390                 395
178       atg aca gca ctg act gga gcg tcc ggt tca gga aaa tcg act ttg ctc      1248
179       Met Thr Ala Leu Thr Gly Ala Ser Gly Ser Gly Lys Ser Thr Leu Leu
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181       aac tgt ctt ggc aca ctt gac aaa cca agt tcc gga cag atc ctt gtc      1296
182       Asn Cys Leu Gly Thr Leu Asp Lys Pro Ser Ser Gly Gln Ile Leu Val
183                   415                 420                 425
184       gag gat gta gac ctt ctg aaa ctc tct acg cgt aag caa cgg tta tac      1344
185       Glu Asp Val Asp Leu Leu Lys Leu Ser Thr Arg Lys Gln Arg Leu Tyr
186               430                 435                 440
187       agg aaa aat acg gtg ggc tat tta ttt caa gat tat gcc ttg att ccc      1392
188       Arg Lys Asn Thr Val Gly Tyr Leu Phe Gln Asp Tyr Ala Leu Ile Pro
189           445                 450                 455
190       gac agg aca gtt aaa ttc aac ctt cag ctt gcg gtg gaa aaa cac aaa      1440
191       Asp Arg Thr Val Lys Phe Asn Leu Gln Leu Ala Val Glu Lys His Lys
192       460                 465                 470                 475
193       tgg cct gaa att cct caa gta ctt cat gct gtt ggt ctt gag tcg ttc      1488
194       Trp Pro Glu Ile Pro Gln Val Leu His Ala Val Gly Leu Glu Ser Phe
195                       480                 485                 490
196       gag gaa aag cca gtt ttt gaa ctc tct ggt ggc gaa caa caa cga act      1536
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RAW SEQUENCE LISTING DATE: 02/26/2002
PATENT APPLICATION: US/09/868,338 TIME: 12:46:49

Input Set : N:\Crf3\02132002\I868338.raw
Output Set: N:\CRF3\02262002\I868338.raw

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201             510                 515                 520
202     gaa cca acc gga gct cta gat tta aca aac agt gag cta gtc ata gaa     1632
203     Glu Pro Thr Gly Ala Leu Asp Leu Thr Asn Ser Glu Leu Val Ile Glu
204         525                 530                 535
205     gca ttg aga gca ctc gcc gac aaa ggc gcc acc gtt gtt gtt gct acg     1680
206     Ala Leu Arg Ala Leu Ala Asp Lys Gly Ala Thr Val Val Val Ala Thr
207     540                 545                 550                 555
208     cac tcg ccc ctc ttc cga gaa tca gcg gat acc att atc aaa cta         1725
209     His Ser Pro Leu Phe Arg Glu Ser Ala Asp Thr Ile Ile Lys Leu
210                     560                 565                 570
211     taggtgcccc aacttttcgg agatctcagt gca atg atg gaa ttc tta aac act     1779
212                                         Met Met Glu Phe Leu Asn Thr
213                                                             575
214     cac cgt ttg att gtt ctc ggg agt ttg tct ttt cta ggg cta ggt ttc     1827
215     His Arg Leu Ile Val Leu Gly Ser Leu Ser Phe Leu Gly Leu Gly Phe
216             580                 585                 590
217     gcg gaa gtc ctg ctg cgt ggc cag tgg tca aca ccg cag ttt ttt gtt     1875
218     Ala Glu Val Leu Leu Arg Gly Gln Trp Ser Thr Pro Gln Phe Phe Val
219         595                 600                 605
220     ttc act ttc ttg caa act ctg ctt ctc gta ttg tgt ttt att cct aaa     1923
221     Phe Thr Phe Leu Gln Thr Leu Leu Leu Val Leu Cys Phe Ile Pro Lys
222     610                 615                 620                 625
223     ctc tcg gtt cct ttt gtg gtg ctt cta agc att gcc caa ctc gcg ctt     1971
224     Leu Ser Val Pro Phe Val Val Leu Leu Ser Ile Ala Gln Leu Ala Leu
225                     630                 635                 640
226     gca tac ctg tgt att cat ggt gaa cct caa agc acc agc cct ttc act     2019
227     Ala Tyr Leu Cys Ile His Gly Glu Pro Gln Ser Thr Ser Pro Phe Thr
228                 645                 650                 655
229     tta att gtt gcc caa atg gcg ttt tcg gga ttg ctc atg ttc aga ggg     2067
230     Leu Ile Val Ala Gln Met Ala Phe Ser Gly Leu Leu Met Phe Arg Gly
231             660                 665                 670
232     caa cgg gtg ctc gct ttt atc tct gca ggt ggg ctc att tgg att ggg     2115
233     Gln Arg Val Leu Ala Phe Ile Ser Ala Gly Gly Leu Ile Trp Ile Gly
234         675                 680                 685
235     acc atc gat cca aca aac ggt gct tgg tct cct cat gtg atg tcc gcg     2163
236     Thr Ile Asp Pro Thr Asn Gly Ala Trp Ser Pro His Val Met Ser Ala
237     690                 695                 700                 705
238     cta gca ctt gcc gta ttc ttt gcg ctg tcg atg gca ctt gga cag gtt     2211
239     Leu Ala Leu Ala Val Phe Phe Ala Leu Ser Met Ala Leu Gly Gln Val
240                     710                 715                 720
241     ctt cga tca aaa gtt gaa caa aga gcc aac ctt gag gag cag gca aaa     2259
242     Leu Arg Ser Lys Val Glu Gln Arg Ala Asn Leu Glu Glu Gln Ala Lys
243                 725                 730                 735
244     att cag aca gaa ctg cgc aga aaa gaa cta agc act cca tct gca tcg     2307
245     Ile Gln Thr Glu Leu Arg Arg Lys Glu Leu Ser Thr Pro Ser Ala Ser
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VERIFICATION SUMMARY DATE: 02/26/2002
PATENT APPLICATION: US/09/868,338 TIME: 12:46:50

Input Set : N:\Crf3\02132002\I868338.raw
Output Set: N:\CRF3\02262002\I868338.raw

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L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2

PCT09

RAW SEQUENCE LISTING DATE: 02/13/2002
PATENT APPLICATION: US/09/868,338 TIME: 07:43:08

Input Set : A:\209861us.ST25.txt
Output Set: N:\CRF3\02132002\I868338.raw

Does Not Comply
Corrected Diskette Needed
P2

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     3       MATSUI, KAZUHIKO
     4       NAKAMATSU, TSUYOSHI
     5       KIMURA, EIICHIRO
     7 <120> TITLE OF INVENTION: ABC TRANSPORTER AND GENE CODING FOR THE SAME
     9 <130> FILE REFERENCE: 209861US-8222-10-0-PCT
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    13 <141> CURRENT FILING DATE: 2001-06-18
    15 <150> PRIOR APPLICATION NUMBER: JP 10/360621
    17 <151> PRIOR FILING DATE: 1998-12-18
    19 <150> PRIOR APPLICATION NUMBER: PCT/JP 99/07079
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    25 <170> SOFTWARE: PatentIn version 3.1
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RAW SEQUENCE LISTING DATE: 02/13/2002
PATENT APPLICATION: US/09/868,338 TIME: 07:43:08

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RAW SEQUENCE LISTING DATE: 02/13/2002
PATENT APPLICATION: US/09/868,338 TIME: 07:43:08

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Output Set: N:\CRF3\02132002\I868338.raw

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197 <222> LOCATION: (1759)..(2367)
199 <223> OTHER INFORMATION:
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204 1               5                  10                  15
206 gga tcg cta acc ggt aac ctc agt gaa cta cgt gca caa act act ttt      96
207 Gly Ser Leu Thr Gly Asn Leu Ser Glu Leu Arg Ala Gln Thr Thr Phe
208             20                  25                  30
210 agt aca tta tgg gat acc cat aaa gaa acc tat aga gtc tcc ata gct     144
211 Ser Thr Leu Trp Asp Thr His Lys Glu Thr Tyr Arg Val Ser Ile Ala
212         35                  40                  45
214 tcc gca gca gga caa gac ttc tac ggg ctt gct gag act cta cgc act     192
215 Ser Ala Ala Gly Gln Asp Phe Tyr Gly Leu Ala Glu Thr Leu Arg Thr
216     50                  55                  60
218 atg gat agg cat ggg gaa att att ttg gca gat cgt caa tgg tta aca     240
219 Met Asp Arg His Gly Glu Ile Ile Leu Ala Asp Arg Gln Trp Leu Thr
220 65                  70                  75                  80
222 gct ccc ctt gat atc ggt gca cca gtc gta tta tca aac aca act ttt     288
223 Ala Pro Leu Asp Ile Gly Ala Pro Val Val Leu Ser Asn Thr Thr Phe
224                 85                  90                  95
226 gcc gtt gat gaa gga cta ctt gcg cca aaa gat cta ccg caa agt gac     336
227 Ala Val Asp Glu Gly Leu Leu Ala Pro Lys Asp Leu Pro Gln Ser Asp
228             100                 105                 110
230 gag atc aca ata ttg cat cct cag ttt ctg gat tcg gcc aaa gag cca     384
231 Glu Ile Thr Ile Leu His Pro Gln Phe Leu Asp Ser Ala Lys Glu Pro
232         115                 120                 125
234 gaa tta ctt ggt ttg ctg gag ttc gaa gca tcc aac tca caa gtg cca     432
235 Glu Leu Leu Gly Leu Leu Glu Phe Glu Ala Ser Asn Ser Gln Val Pro
236     130                 135                 140
238 atg cca aag atc caa agc att cca tat gat agc gaa gac tca acc aac     480
239 Met Pro Lys Ile Gln Ser Ile Pro Tyr Asp Ser Glu Asp Ser Thr Asn
240 145                 150                 155                 160
242 ccc atg tct gaa gtt ttt acc tac aac att aac ctg gat agt gca gta     528
243 Pro Met Ser Glu Val Phe Thr Tyr Asn Ile Asn Leu Asp Ser Ala Val
244                 165                 170                 175
246 aga aac cca atc gta gtt atc ctt ccc gca ggc tta gag ctt tta agt     576
247 Arg Asn Pro Ile Val Val Ile Leu Pro Ala Gly Leu Glu Leu Leu Ser
248             180                 185                 190
250 gat caa aat ttg tcg gct cga ctc aca cag aat agt ctg ctg ata aaa     624
251 Asp Gln Asn Leu Ser Ala Arg Leu Thr Gln Asn Ser Leu Leu Ile Lys
252         195                 200                 205
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RAW SEQUENCE LISTING DATE: 02/13/2002
PATENT APPLICATION: US/09/868,338 TIME: 07:43:08

Input Set : A:\20986lus.ST25.txt
Output Set: N:\CRF3\02132002\I868338.raw

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254 gac cag act ggt gtg aac gct ctt cta tcc tca gag gat tca cgc aat      672
255 Asp Gln Thr Gly Val Asn Ala Leu Leu Ser Ser Glu Asp Ser Arg Asn
256     210                 215                 220
258 tat gtg gga gct gca tcc ccg atg att gac acg tgg gaa gaa agc gtt      720
259 Tyr Val Gly Ala Ala Ser Pro Met Ile Asp Thr Trp Glu Glu Ser Val
260 225                 230                 235                 240
262 gtt cgg ttg aag gaa gcg aac caa ata atc gcc ttc aac gct ttc att      768
263 Val Arg Leu Lys Glu Ala Asn Gln Ile Ile Ala Phe Asn Ala Phe Ile
264                 245                 250                 255
266 gca ttg ttc ctc acg acg act ctt gtt cta gca tac tgc act ggt att      816
267 Ala Leu Phe Leu Thr Thr Thr Leu Val Leu Ala Tyr Cys Thr Gly Ile
268             260                 265                 270
270 tca ttt aag aaa tca aag aag act atg ggt agc gca tct act agg aaa      864
271 Ser Phe Lys Lys Ser Lys Lys Thr Met Gly Ser Ala Ser Thr Arg Lys
272         275                 280                 285
274 tca tcc att aag agc tcg att aca gct gct aat tgt aga agt aat ttt      912
275 Ser Ser Ile Lys Ser Ser Ile Thr Ala Ala Asn Cys Arg Ser Asn Phe
276     290                 295                 300
278 cgc ttc aat tcc gtg cgt ctg gct cgc gaa ccg cta ttt cga gcg atc      960
279 Arg Phe Asn Ser Val Arg Leu Ala Arg Glu Pro Leu Phe Arg Ala Ile
280 305                 310                 315                 320
282 tgc agc aat agc ttc aga tgc tcc ctc agc cag ata ctt aga aca tct     1008
283 Cys Ser Asn Ser Phe Arg Cys Ser Leu Ser Gln Ile Leu Arg Thr Ser
284                 325                 330                 335
286 caa ttc tat acc tcc atc act gcc gtt ggt ttt agg aat ctt aat aat     1056
287 Gln Phe Tyr Thr Ser Ile Thr Ala Val Gly Phe Arg Asn Leu Asn Asn
288             340                 345                 350
290 cgg ttg gac ttc act ttc att ttt cag ttc gat gaa gct tcc ttt         1101
291 Arg Leu Asp Phe Thr Phe Ile Phe Gln Phe Asp Glu Ala Ser Phe
292         355                 360                 365
294 tgaaaagagc acaca atg ata gaa atc aat gac ctc aag aaa tct ttt ggc    1152
295                 Met Ile Glu Ile Asn Asp Leu Lys Lys Ser Phe Gly
296                     370                 375
298 gtt cgg atc tta tgg caa ggt ctc agt cat aag ttt tta cca gga aca     1200
299 Val Arg Ile Leu Trp Gln Gly Leu Ser His Lys Phe Leu Pro Gly Thr
300 380                 385                 390                 395
302 atg aca gca ctg act gga gcg tcc ggt tca gga aaa tcg act ttg ctc     1248
303 Met Thr Ala Leu Thr Gly Ala Ser Gly Ser Gly Lys Ser Thr Leu Leu
304                 400                 405                 410
306 aac tgt ctt ggc aca ctt gac aaa cca agt tcc gga cag atc ctt gtc     1296
307 Asn Cys Leu Gly Thr Leu Asp Lys Pro Ser Ser Gly Gln Ile Leu Val
308             415                 420                 425
310 gag gat gta gac ctt ctg aaa ctc tct acg cgt aag caa cgg tta tac     1344
311 Glu Asp Val Asp Leu Leu Lys Leu Ser Thr Arg Lys Gln Arg Leu Tyr
312         430                 435                 440
314 agg aaa aat acg gtg ggc tat tta ttt caa gat tat gcc ttg att ccc     1392
315 Arg Lys Asn Thr Val Gly Tyr Leu Phe Gln Asp Tyr Ala Leu Ile Pro
316     445                 450                 455
318 gac agg aca gtt aaa ttc aac ctt cag ctt gcg gtg gaa aaa cac aaa     1440
```

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319 Asp Arg Thr Val Lys Phe Asn Leu Gln Leu Ala Val Glu Lys His Lys
320 460                 465                 470                 475
322 tgg cct gaa att cct caa gta ctt cat gct gtt ggt ctt gag tcg ttc    1488
323 Trp Pro Glu Ile Pro Gln Val Leu His Ala Val Gly Leu Glu Ser Phe
324                 480                 485                 490
326 gag gaa aag cca gtt ttt gaa ctc tct ggt ggc gaa caa caa cga act    1536
327 Glu Glu Lys Pro Val Phe Glu Leu Ser Gly Gly Glu Gln Gln Arg Thr
328             495                 500                 505
330 gcg ttg gcc cgg gta ctg ctc aaa aat ccc cga ata att ctg gct gat    1584
331 Ala Leu Ala Arg Val Leu Leu Lys Asn Pro Arg Ile Ile Leu Ala Asp
332         510                 515                 520
334 gaa cca acc gga gct cta gat tta aca aac agt gag cta gtc ata gaa    1632
335 Glu Pro Thr Gly Ala Leu Asp Leu Thr Asn Ser Glu Leu Val Ile Glu
336     525                 530                 535
338 gca ttg aga gca ctc gcc gac aaa ggc gcc acc gtt gtt gtt gct acg    1680
339 Ala Leu Arg Ala Leu Ala Asp Lys Gly Ala Thr Val Val Val Ala Thr
340 540                 545                 550                 555
342 cac tcg ccc ctc ttc cga gaa tca gcg gat acc att atc aaa cta       1725
343 His Ser Pro Leu Phe Arg Glu Ser Ala Asp Thr Ile Ile Lys Leu
344                 560                 565                 570
346 taggtgcccc aacttttcgg agatctcagt gca atg atg gaa ttc tta aac act    1779
347                                      Met Met Glu Phe Leu Asn Thr
348                                                      575
350 cac cgt ttg att gtt ctc ggg agt ttg tct ttt cta ggg cta ggt ttc    1827
351 His Arg Leu Ile Val Leu Gly Ser Leu Ser Phe Leu Gly Leu Gly Phe
352         580                 585                 590
354 gcg gaa gtc ctg ctg cgt ggc cag tgg tca aca ccg cag ttt ttt gtt    1875
355 Ala Glu Val Leu Leu Arg Gly Gln Trp Ser Thr Pro Gln Phe Phe Val
356     595                 600                 605
358 ttc act ttc ttg caa act ctg ctt ctc gta ttg tgt ttt att cct aaa    1923
359 Phe Thr Phe Leu Gln Thr Leu Leu Leu Val Leu Cys Phe Ile Pro Lys
360 610                 615                 620                 625
362 ctc tcg gtt cct ttt gtg gtg ctt cta agc att gcc caa ctc gcg ctt    1971
363 Leu Ser Val Pro Phe Val Val Leu Leu Ser Ile Ala Gln Leu Ala Leu
364                 630                 635                 640
366 gca tac ctg tgt att cat ggt gaa cct caa agc acc agc cct ttc act    2019
367 Ala Tyr Leu Cys Ile His Gly Glu Pro Gln Ser Thr Ser Pro Phe Thr
368             645                 650                 655
370 tta att gtt gcc caa atg gcg ttt tcg gga ttg ctc atg ttc aga ggg    2067
371 Leu Ile Val Ala Gln Met Ala Phe Ser Gly Leu Leu Met Phe Arg Gly
372         660                 665                 670
374 caa cgg gtg ctc gct ttt atc tct gca ggt ggg ctc att tgg att ggg    2115
375 Gln Arg Val Leu Ala Phe Ile Ser Ala Gly Gly Leu Ile Trp Ile Gly
376     675                 680                 685
378 acc atc gat cca aca aac ggt gct tgg tct cct cat gtg atg tcc gcg    2163
379 Thr Ile Asp Pro Thr Asn Gly Ala Trp Ser Pro His Val Met Ser Ala
380 690                 695                 700                 705
382 cta gca ctt gcc gta ttc ttt gcg ctg tcg atg gca ctt gga cag gtt    2211
383 Leu Ala Leu Ala Val Phe Phe Ala Leu Ser Met Ala Leu Gly Gln Val
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VERIFICATION SUMMARY DATE: 02/13/2002
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Input Set : A:\20986lus.ST25.txt
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L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:64 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:181 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:7